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Listing first 45 summaries
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Perfect score:
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    103943 segs, 16242309 residues
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1747
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241 FSVPFTPYHVMRNVRIASRLGSWKQYQCTQVVINSFYIVTRPLAFLNSVINPVFYFLLGD

PNYNLIYSMCLTLLGFLIPLFVMCFFYYKIALFLKQRNRQVATALPLEKPLNLVIMAVVI

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YLIIKYPFREHLLOKKEFAILISLAIWVLVTLELLPILPLINPVITDNGTTCNDFASSGD

Qy 1 MLGIMAWNATCKNWLAAEAALEKYYLSIFYGIEFVVGVLGNTIVVYGYIFSLKNWNSSNI 60	RESULT 1 US-09-765-034-2 US-09-765-034-2 Sequence 2, Application US/09765034 Patent No. US20020137887A1 GENERAL INFORMATION: APPLICANT: Hedrick, Joseph A. APPLICANT: Hedrick, Joseph A. APPLICANT: Gustafson, Eric L. TITLE OF INVENTION: Adenosine Receptor FILE REFERENCE: CN01084 CURRENT FILING DATE: 2001-01-17 NUMBER OF SEQ ID NOS: 4 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 2 LENGTH: 334 TYPE: PRT ORGANISM: homo sapiens US-09-765-034-2 Query Match Best Local Similarity 100.0%; Score 1747; DB 10; Best Local Similarity 100.0%; Pred. No. 4.3e-142; Matches 334; Conservative 0; Mismatches 0;	ALIGNMENTS	20 326.5 21 325 22 321.5 23 325 24 321.5 26 31.5 27 31.7 30 317.5 31 312.5 36 312.5 37 312.5 38 312.5 39 311.5 31 312.5
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                                          301 HFRDMLMNQLRHNFKSLTSFSRWAHELLLSFREK 334
                                                             301 HFRDMLMNQLRHNFKSLTSFSRWAHELLLSFREK 334
                                                                                                                                       Sequence 4, Application US/09943798
Patent No. US20020065215A1
GENERAL INFORMATION:
TITLE OF INVENTION: Polypeptide
FILE REFERENCE: QC1021
CURRENT APPLICATION NUMBER: US/09/943,798
CURRENT FILING DATE: 2001-08-31
NUMBER OF SEQ ID NOS: 4
SEQ ID NO 4
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Glaxo Group Limited
TITLE OF INVENTION: Polypeptide
FILE REFERENCE: QG1021
CURRENT APPLICATION NUMBER: US/09/943,798
CURRENT FILING DATE: 2001-08-31
NUMBER OF SEQ ID NOS: 4
SCOTTANE: FastSEQ for Windows Version 3.0
SEQ ID NO 2
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Patent No. US20020065215A1
GENERAL INFORMATION:
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ilarity 35.9%;
Conservative 63
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US-09-943-798-4
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ORGANISM: Homo sapiens
US-09-943-798-2
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     Length 276;
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Query Match 22.6%; Score 395.5; DB 10; Length Best Local Similarity 35.1%; Pred. No. 7e-27; Matches 92; Conservative 54; Mismatches 109; Indels
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Facture No. US20010039037A1

GENERAL INFORMATION:
APPLICANT: Pfizer Inc.
TITLE OF INVENTION: NO. US20010039037A1e1 Polypeptide
FILE REPRENCE: PC10914AAAA

CURRENT APPLICATION NUMBER: US/09/826,791

CURRENT FILING DATE: 2001-04-05

PRIOR APPLICATION NUMBER: 60/198,367

PRIOR FILING DATE: 2000-04-15

SOFTWARE: PALCHIN VOR: 6

SOFTWARE: PALCHIN VOR: 2.0
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; ORGANISM: Homo sapiens
US-09-826-791-2
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Best Local Similarity
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                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Pfizer Inc
TITLE OF INVENTION: NO. US20010039037A1el Polypeptide
FILE REFERENCE: PC10914ADAM
                                                                                                                                                                                                                                                                                     Sequence 6, Application US/09826791 Patent No. US20010039037A1
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APPLICANT: Xiao, Yonghong
TITLE OF INVENTION: Regulation of Human CysLT2-Like
TITLE OF INVENTION: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
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Patent No. US200
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                                                            CURRENT APPLICATION NUMBER: US/09/826,791
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 0008504.3
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: 60/198,367
PRIOR APPLICATION NUMBER: 60/198,367
PRIOR FILING DATE: 2000-04-19
SOFTWARE: PatentIn Ver. 2.0 EQ ID NO 6
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PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: 60/254,876
PRIOR FILING DATE: 2000-12-13
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CURRENT FILING DATE: 2001-04-09
                                               NUMBER OF SEQ ID NOS: 6
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US20020155528A1
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; LENGTH: 346
; TYPE: PRT
; ORGANISM: Homo s
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US-09-826-791-6
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APPLICANT: MUTPHY, Andrew, et al.
TITLE OF INVENTION: No. US20020150901A1el Nucleic
FILE REFERENCE: REG 771A
CURRENT APPLICATION NUMBER: US/09/866,230
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.0 SEQ ID NO 7
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                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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Best Local Similarity
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PRIOR FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 9
                                      183 YNLIYSMCLTL-----
                                                                                                             124 IKYPFR-EHLLQKKEFAILISLAIWVLVTLELLPILPLINPVITDNGTTCNDFASSGDPN 182
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                                                                                                                                                                                       LSVSDLAFLCTLPMLIRSYANG-NWIYGDVLCISNRYVLHANLYTSILFLTFISIDRYLI 123
                                                                          MVHPFRLLHVTSIRSAWILCGI-IWILIMASSIMLL
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CLELNLYKIAKLQTMNYIALVVGCLLPFFTLSICYLLIIRVLLKVEVPESGLR 239
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                                                                                                                                                                                                                                                                                                                             22.0%; Score 384; DB 10; 30.4%; Pred. No. 8.6e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . 59; Mismatches 125;
                                                                                                                                                                                                                                                                                                              59;
                                                                                                                                                                                                                                                                                                              Mismatches
                                    -LGFLIPLFVMCFFYYK-IALFLKQRNRQVATA 224
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                                                                                                                                                                                                                                                                                                              125;
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Sequence 8, Application US/09866230
Patent No. US20020150901A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: No. US20020150901A1e1 Nucleic Acids, Polypeptides, Methods of Ma
FILE REFERENCE: REG 771A
CURRENT APPLICATION NUMBER: US/09/866,230
CURRENT FILING DATE: 2001-05-25
PRIOR PAPLICATION NUMBER: 60/207,725
PRIOR FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68 SDLAFLCTLPMLIRSYAN-GNWIYGDVLCISNRYVLHANLYTSILFLTFISIDRYLIIKY 126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         184 NLIYSMCLTLLGFLIP--LFVMCFFYYKIALFLKQRNRQVATALPLEKPLNLVIMAVVIF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              127 PFREHLLQKKEFAILISLAIWVLVTLELLPILPLINPVITDNGTTCNDFASSGD---PNY
                                                                                                                                                                                                                                                                               8 NATCKNWLAAEAALEKYYLSIFYGIEFVVGVLGNTIVVYGYIFSLKNWNSSNIYLFNLSV
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                                                                                                                                                                                    Query Match 20.3%; Score 354.5; DB 9; Best Local Similarity 28.3%; Pred. No. 2.8e-23; Matches 91; Conservative 62; Mismatches 152;
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             301 NFRKRLSTFRKHSLSSVTYVPR 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                301 HFRDMLMNQLRHNFKSLTSFSR 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.0
                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
US-09-828-478-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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  NUMBE: FABLE
SOFTWARE: FABLE
SEQ ID NO 5
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US-09-866-230-8
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TITLE OF INVENTION: No. US20020150901Alel Nucleic Acids, Polypeptides, Methods of Mak
TITLE OF INVENTION: No. US20020150901Alel Nucleic Acids, Polypeptides, Methods of Mak
CHERDY APPLICATION NUMBER: US/09/866,230
CURRENT APPLICATION NUMBER: 60/207,725
PRIOR FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 9
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                            124 IKYPFR-EHLLQKKEFAILISLAIWVLVTLELLPILPLINPVITDNGTTCNDFASSGDPN 182
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  183 YNLIYSMCLTL--------LGFLIPLFVMCFFYYK-IALFLKQNATA
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Sequence 5, Application US/09828478
Sequence 5, Application US/09828478
Patent No. US20020155528A1
GENERAL INFORMATION: Regulation of Human CyeLT2-Like GPCR
TITLE OF INVENTION: Regulation of Human CyeLT2-Like GPCR
TITLE OF INVENTION: Protein
FILE REFERENCE: 04974.00458
CURRENT PELLING DATE: 2001-04-09
PRIOR APPLICATION NUMBER: 60/195,196
PRIOR PLILING DATE: 2000-04-07
PRIOR PLILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 22.0%; Score 384; DB 10; Best Local Similarity 30.4%; Pred. No. 8.6e-26; Matches 102; Conservative 59; Mismatches 125;
                                                                                                                     295 AANACFNPLLYYFAGENFKDRLKSALRKGHPQKAKT 330
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                                                                                                                                                                                                                                                       Sequence 9, Application US/09866230 Patent No. US20020150901A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.0
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US-09-866-230-9
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                                                                                                                                                                                                         RESULT 8
US-09-866-230-9
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US-09-828-478-4
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APPLICANT: Li, Yi
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human G-Protein Coupled Refere Reference: 1488.1220003
CURRENT APPLICATION NUMBER: US/09/827,937A
CURRENT FILING DATE: 2001-04-09
PRIOR APPLICATION NUMBER: 08/852,824
PRIOR FILING DATE: 1997-05-07
NUMBER OF SEQ ID NOS: 18
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GENERAL INFORMATION:
APPLICANT: Xiao, Yonghong
APPLICANT: Xiao, Yonghong
TITLE OF INVENTION: Regulation of Human CysLT2-Like GPCR
TITLE OF INVENTION: Protein
FILE REFERENCE: 04974.00458
CURRENT APPLICATION NUMBER: US/09/828,478
                                                                                                                                      Sequence 4, Application US/09828478 Patent No. US20020155528A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 19.8%; Score 346; DB 10; Best Local Similarity 28.7%; Pred. No. 1.5e-22; Matches 95; Conservative 69; Mismatches 135;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 LAAEAALEKYYLSIFYGIEFVVGVLGNTIVVYGYIFSLKNWNSSNIYLFNLSVSDLAFLC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILLGACFIGYVLPLIIIKICYSQICCKLFRTAKQ----NPLTEKSGVNKKALNTIILIIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TLL-----GFLIPLFVMCFFYYKIALFLKQRNRQVATALPL-----EKPLNLVIMAVV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HFRDMLMNQLRHNFKSLTSFSR 322
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                                                                                                                                                                                                                                                                              FACKGYKRKVMRMLKRQVSVSISSAVKSAPE
                                                                                                                                                                                                                                                                                                                                                                   VFVLCFTPYHVAIIQHMIKKLRFSNFLECSQ--RHSFQISLHFTVCLMNFNCCMDPFIYF
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GENERAL INFORMATION:

APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.

APPLICANT: Cheng, Muzong
TITLE OF INVENTION: GPCR DIAGNOSTIC FOR BRAIN CANCER
FILE REFERENCE: PC-0042 CIP
CURRENT APPLICATION NUMBER: US/09/848,889
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PERL Program
SEQ ID NO 12
LENGTH: 339
TYPE: PT
ORGANISM: Homo sapiens
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PRIOR APPLICATION NUMBER: 60/195,196
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: 60/254,876
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.
SEQ ID NO 4
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Patent No. US20020025555A1
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                                                                                                                                  Query Match
                                                                                                               Best
                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc feat
OTHER INFORMATION:
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ORGANISM: Homo sapiens
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                                                                                                               Local
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                                     18 EAALEKYYLSIFYGIEFVVGVLGNTIVVYGYIFSLKNWNSSNIYLFNLSVSDLAFLCTLP 77
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ETPLENMLFASFYLLDFILALVGNTLALWLFIRDHKSGTPANVFLMHLAVADLSCVLVLP
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                                                                                                            Similarity
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                                                                                         Conservative
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                                                                                     19.3%; Score 338; DB 10; 27.5%; Pred. No. 7.1e-22; tive 65; Mismatches 126
                                                                                                                             Length 339;
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                    EPAILISLAIWVLVTLELLPILPLINPVITDNGTTCNDFASSGDPNYNLIYSMCLTLLGF 196
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19.3%; Score 338; DB 10; Length 339;
Best Local Similarity 27.5%; Pred. No. 7.1e-22;
Matches 84; Conservative 65; Mismatches 126; Indels 3
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Patent No. US2002065201A1
GENERAL INFORMATION:
TITLE OF INVENTION: ASSAY
FILE REFRENCE: P79011
CURRENT APPLICATION NUMBER: US/09/788,133
CURRENT FILING DATE: 2001-02-20
NUMBER OF SEQ ID NOS: 2
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
CRGANISM: homo sapiens
US-09-788-133-2
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US-09-788-133-2
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Sequence 6, Application US/09828478; Patent No. US20020155528A1; GENERAL INFORMATION:

US-09-828-478-6

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230 TFPFITTVTCYLLIIRSLRQ------GLRVEKRLKTKAVRMIAIVLAIFLVCFVPYHVN 282
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                                                                                                                                                                                                                                                                                                                                  ; DB 9; Length 367; 7.7e-22;
TITLE OF INVENTION: Regulation of Human CysLT2-Like GPCR
TITLE OF INVENTION: Protein
FILE REPERENCE: 04974-00458
CURRENT APPLICATION NUMBER: US/09/828,478
CURRENT FILING DATE: 2001-04-09
PRIOR APPLICATION NUMBER: 60/195,196
PRIOR PILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 16
SEQ ID NOS: 16
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                        65; Mismatches 126; Indels
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Job time : 12 secs
                                                                                                                                                                                                                                                                                                                                                                        84; Conservative
                                                                                                                                                                                                                                 LENGTH: 367
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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